

053-10280

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/067,988

DATE: 02/28/2002
TIME: 12:34:11

Input Set : A:\2314-210-II.ST25.txt
Output Set: N:\CRF3\02282002\J067988.raw

3 <110> APPLICANT: Bulaj, Grzegorz
 5 <120> TITLE OF INVENTION: Methods for Refolding Conformationally Constrained Peptides
 7 <130> FILE REFERENCE: 2314-210-II
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/067,988
 C--> 9 <141> CURRENT FILING DATE: 2002-02-08
 9 <150> PRIOR APPLICATION NUMBER: US 60/267,192
 10 <151> PRIOR FILING DATE: 2001-02-08
 12 <160> NUMBER OF SEQ ID NOS: 7
 14 <170> SOFTWARE: PatentIn version 3.0
 16 <210> SEQ ID NO: 1
 17 <211> LENGTH: 29
 18 <212> TYPE: PRT
 19 <213> ORGANISM: Conus purpurascens
 21 <400> SEQUENCE: 1
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 30 <211> LENGTH: 31
 31 <212> TYPE: PRT
 32 <213> ORGANISM: Conus striatus
 34 <400> SEQUENCE: 2
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 44 <212> TYPE: PRT
 45 <213> ORGANISM: Conus textile
 47 <400> SEQUENCE: 3
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 57 <212> TYPE: PRT
 58 <213> ORGANISM: Conus gloriamaris
 60 <400> SEQUENCE: 4
 62 Val Lys Pro Cys Arg Lys Glu Gly Gln Leu Cys Asp Pro Ile Phe Gln
 63 1 5 10 15

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65 Asn Cys Cys Arg Gly Trp Asn Cys Val Leu Phe Cys Val
66 20 25
68 <210> SEQ ID NO: 5
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70 <212> TYPE: PRT
71 <213> ORGANISM: Conus marmoreus
73 <400> SEQUENCE: 5
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78 Val Tyr Cys Cys Pro Gly Leu Ile Cys Gly Phe Val Cys Val
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81 <210> SEQ ID NO: 6
82 <211> LENGTH: 27
83 <212> TYPE: PRT
84 <213> ORGANISM: Conus gloriamaris
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89 1 5 10 15
91 Ile Cys Thr Phe Arg Gly Cys Gly Ala Val Asn
92 20 25
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96 <212> TYPE: PRT
97 <213> ORGANISM: Conus textile
99 <220> FEATURE:
100 <221> NAME/KEY: PEPTIDE
101 <222> LOCATION: (1)..(27)
102 <223> OTHER INFORMATION: Xaa is gamma-carboxy-Glu
105 <400> SEQUENCE: 7
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108 1 5 10 15
110 Ile Cys Thr Phe Arg Gly Cys Gly Ala Val Asn
111 20 25

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/067,988

DATE: 02/28/2002
TIME: 12:34:12

Input Set : A:\2314-210-II.ST25.txt
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L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7